WHAT IS CLAIMED IS:

1. An oligopeptide that comprises a sequence of amino acids that is recognized and selectively proteolytically cleaved by free prostate specific antigen.

2. The oligopeptide according to Claim 1 wherein the sequence of amino acids is

a) AsnLysIleSerTyrGln|Ser (SEQ.ID.NO.: 13),

b) LysIleSerTyrGln|Ser (SEQ.ID.NØ.: 14),

c) GlyGluAsnGlyValGlnLysAspValSerGlnXaaSerIleTyr|SerGlnThrGlu (SEQ.ID.NO.: 15),

d) GlyLysGlyIleSerSerGlnTyr|SerAsnThrGluGluArgLeu (SEQ.ID.NO.: 2),

e) AsnLysIleSerTyrTyr|Ser

(SEQ.ID.NO.: 127),

f) AsnLysAlaSerTyrGln|Ser

(SEQ.ID.NO.: 128),

g) SerTyrGln|SerSer

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(SEQ.HD.NO.: 129);

h) LysTyrGln|SerSer

(SEQ.ID.NO.: 140); or

i) hArgTyrGln|SerSer

(SEQ.ID.NO.: 141);

wherein hArg is homoarginine and Xaa is any natural amino acid.

3. The oligopeptide according to Claim 2 wherein the sequence of amino acids is

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a) AsnLysIleSerTyrGln|SerSer (SEQ.ID.NO./16),

b) AsnLysIleSerTyrGln|SerAla (SEQ.ID.No.: 130),

c) AsnLysIleSerTyrGln|SerSerSer (SEQ.ID.NO.: 17),

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d) AlaAsnLysIleSerTyrGln|SerSerSer (SEQ.ID.NO.: 18),

e) LysIleSerTyrGln|SerSerSerThrGlu (SEQ.ID.NO.: 19),

f) GlyGluAsnGlyValGlnLysAspValSerGlnArgSerIleTyr|SerGlnThrGlu (SEQ.ID.NO.: 4),

g) GlyGluAsnGlyValGlnLysAspValSerGlnSerSerIleTyr|SerGlnThrGlu (SEQ.ID.NO.: 5),

h) AlaAsnLysIleSerTyrTyr|Ser (SEQ.ID.NO.: 131),

i) AlaAsnLysAlaSerTyrGln|Ser / (SEQ.ID.NO.: 132),

j) SerTyrGln|SerSerThr (SEQ.ID.NO.: 133),

k) SerTyrGln|SerSerSer / (SEQ.ID.NO.: 134),

25 l) LysTyrGln|SerSerSer (SEQ.ID.NO.: 142),

m) hArgTyrGln|SerSerSer (SEQ.ID.NO.: 143), or

n) SerTyrGln|SerSerLeu (SEQ.ID.NO.: 135).

4. The oligopeptide according to Claim 2 wherein the amino acid sequence is

	a) AsnLysIleSerTyrGln SerSerSerThr (SEQ.ID.NO.: 10),
	b) AlaAsnLysIleSerTyrGln SerAla (SEQ.ID.NO.: 136),
5	c) AsnLysIleSerTyrGln SerSerSerThrGlu (SEQ.ID.NO.:3),
	d) AlaAsnLysIleSerTyrGln SerSerSerThrGlu / (SEQ.ID.NO.: 11),
10	e) GlyGluAsnGlyValGlnLysAspValSerGlnArgSerIleTyr SerGlnThrGlu (SEQ.ID.NO.: 4),
	f) AlaAsnLysIleSerTyrTyr SerSer (SEQ.ID.NO.: 137),
15	g) AlaAsnLysIleSerTyrTyr SerAla (SEQ.ID.NO.: 138),
	h) AlaAsnLysAlaSerTyrGln SerAla /(SEQ.ID.NO.: 139), or
	i) AlaSerTyrGln SerSerLeu (SEØ/ID.NO.: 94).
20	5. The oligopeptide according to Claim 2 wherein the amino acid sequence is
	a) GlyArgLysAlaAsnLysIleSerPyrGln SerSerSerThrGluGluArgArg
25	LeuHisTyr GlyGluAsnGly (SEQ.ID.NO.: 6).
	6. The oligopeptide according to Claim 1 which is
	selected from:
30	AsnArgIleSerTyrGln Ser (SEQ.ID.NO.: 21),
	AsnLysValSerTyrGln Ser (SEQ.ID.NO.: 22),
	AsnLysMetSerTyrGln SerSer (SEQ.ID.NO.: 23),
	AsnLysLeuSerTyrGln/SerSer (SEQ.ID.NO.: 24),
	AsnLysIleThrTyrGln SerSerSer (SEQ.ID.NO.: 25),

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AsnLysIleSerPheGln|SerSerSer
                                      (SEQ.ID.NO.
                                                   26),
     AsnLysIleSerTrpGln|SerSerSerThr
                                         (SEQ.ID.NO.: 27),
     AsnLysIleSerTyrAsn|SerSerSerThr
                                         (SEO.ID.NO.: 28),
     AsnLysIleSerTyrGln|ThrSerSerThr
                                         (SEQ.ID.NO.: 29),
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     AsnLysIleSerTyrGln|Ser
                                (SEQ.ID.NO.: 30).
                                  (SEQ.ID.NO £ 31),
     GlnLysIleSerTyrGln|SerSer
     AsnArgIleThrTyrGln|SerSerSer
                                      (SEQ.ID.NO.: 32),
     AsnArgIleSerPheGln|SerSerSerThr
                                         (SEQ.ID.NO.: 33),
     AsnArgIleSerTrpGln|SerSerSerThr
                                         (SEQ.ID.NO.: 35),
10
                                         (SEQ.ID.NO.: 36),
     AsnArgIleSerTyrGln|ThrSerSerThr
     AsnLysIleThrTyrGln|ThrSerSerThr
                                         (SEQ.ID.NO.: 37),
                                          (SEQ.ID.NO.: 38),
     AsnLysLeuSerTyrGln|ThrSerSerThr
     GlnLysLeuSerTyrGln|SerSerSerThr
                                          SEO.ID.NO.: 39),
     AsnArgLeuSerTyrGln|ThrSerSerThr
                                          (SEQ.ID.NO.: 40),
15
                                          (SEQ.ID.NO.: 41),
     AsnLysValSerPheGln|SerSerSerThr
                                          (SEO.ID.NO.: 42),
     AsnArgValSerTrpGln|SerSerSerThr
                                         (SEQ.ID.NO.: 43),
     GlnLysValSerTyrGln|SerSerSerThr
     GlnLysIleSerTyrGln|ThrSerSerThr
                                         (SEQ.ID.NO.: 34), or
     AsnLysIleSerTyrGln|SerSerSerThr
                                         (SEQ.ID.NO.: 44).
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                     The oligopeptide according to Claim 1 which is
                7.
    AlaAsnLysIleSerTyrGln|SerSerSerThrGlu-amide
                                                  (SEQ.ID.NO.: 11)
    Ac-AlaAsnLysIleSerTyrGln|SerSerSerThrLeu
                                                 (SEQ.ID.NO.: 70)
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    Ac-AlaAsnLysIleSerTyrGln|SerSerThrGlu-amide (SEQ.ID.NO.: 11)
    Ac-AlaAsnLysIleSerTyrCln|SerSerSerThrLeu-amide (SEQ.ID.NO.: 70)
    Ac-AlaAsnLysIleSerTyrGlnSerAlaSerThrGlu-amide (SEQ.ID.NO.: 73)
    Ac-AlaAsnLysIleSerTyrGln|\$erSerLysThrGlu-amide (SEQ.ID.NO.: 74)
30
    Ac-AlaAsnLysIleSerTyrGln/SerSerThrGlu-amide
                                                   (SEQ.ID.NO.: 75)
    Ac-AlaAsnLysIleSerTyrGln|SerSerGlnThrGlu-amide (SEQ.ID.NO.: 78)
    Ac-AlaAsnLysIleSerTyrG/in|SerAlaLysThrGlu-amide (SEQ.ID.NO.:79)
    Ac-AlaAsnLysIleSerTyrGln|SerThrGlu-amide
                                                (SEQ.ID.NO.: 81)
    Ac-AlaAsnLysSerTyrGl/n|SerSerThrGlu-amide
                                                 (SEQ.ID.NO.: 82)
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Ac-AlaAsnLysAlaSerTyrGln|SerAlaSerThrGlu-amide (SEQ.ID.NO.: 84) Ac-AlaAsnGluIleSerTyrGln|SerAlaSerThrGlu-amide (SEQ.ID.NO.: 85) Ac-AsnLysIleSerTyrGln|SerSer-amide (SEQ/ID.NO.: 16) (SEQ.ID/NO.: 86) Ac-LysIleSerTyrGln|SerSer-amide Ac-SerTyrGln|SerSerThrGlu-amide (SEQ.ID.NO.: 87) Ac-AlaSerTyrGln|SerSerThrGlu-amide (SEQ.ID.NO.: 89) Ac-AlaAsnLysIleSerTyrTyr|SerSerSerThr|Glu-amide (SEQ.ID.NO.: 92) Ac-AlaAsnLysIleSerTyrTyr|SerAlaSerThrGlu-amide (SEQ.ID.NO.: 93) (SEQ.ID.NO.: 94) Ac-AlaSerTyrGln|SerSerLeu-amide

Ac-AlaAsnSerTyrGln|SerSerSerThrGlu/-amide (SEQ.ID.NO.: 95) Ac-AlaSerTyrGln|SerSerSerThrGlu-amide (SEQ.ID.NO.: 96) Ac-SerTyrGln|SerSerSerThrGlu-amide (SEQ.ID.NO.: 97) or Ac-AlaAsnLysAlaSerTyrGln|SerAlaSerCys-amide (SEQ.ID.NO.: 98).

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- An assay, for determining proteolytic activity of free prostate specific antigen in a sample, comprising the steps of:
 - reacting a substrate, wherein the substrate is an (a), oligopeptide that comprises a sequence of amino acids that is recognized and selectively proteolytically cleaved by free prostate specific antigen, with the sample; and
 - detecting whether the substrate has been cleaved. (b),

- The assay according to Claim 8 wherein the step of 9. detecting whether the substrate has been cleaved comprises analyzing the assay mixture by high performance liquid chromatography.
- An askay for identifying compounds which inhibit the 10. 30 proteolytic activity of prostate specific antigen, comprising:
 - reacting a substrate, wherein the substrate (a), comprises a sequence of amino acids that is recognized and selectively proteolytically cleaved by free prostate specific antigen, with free prostate

specific antigen in the presence of a test substance; and

- (b), detecting whether the substrate has been cleaved, in which the ability of the test substance to inhibit proteolytic activity of prostate specific antigen is indicated by a decrease in the cleavage of the substrate in the absence of the test substance.
- 11. The assay according to Claim 10 wherein the step of detecting whether the substrate has been cleaved comprises analyzing the assay mixture by high performance liquid chromatography.
- prostate cancer which comprises a cytotoxic agent attached to a oligopeptide, wherein the oligopeptide comprises a sequence of amino acids that is recognized and selectively proteolytically cleaved by free prostate specific antigen, wherein the means of attachment is a covalent bond or a chemical linker.

The conjugate according to Claim 12 wherein the cytotoxic agent is a member of a class of cytotoxic agents selected from the following classes:

- a) anthracycline family of drugs,
- b) the vinca alkaloid drugs,
- c) the mitomycins,
- d) the bleomycins,
- e) the cytotoxic nucleosides,
- f) the pteridine family of drugs,
- g) diynenes,
- h) estramustine,
- i) cyclophosphamide, and
- h) the podophyllotoxins.

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The conjugate according to Claim 12 wherein the cytotoxic agent is selected from the following cytotoxic agents:

- a) doxorubicin,
- b) carminomycin,
- c) daunorubicin,
- d) aminopterin,
- e) methotrexate,
- f) methopterin,
- g) dichloro-methotrexate,
- h) mitomycin C,

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- i) porfiromycin,
- j) 5-fluorouracil,
- k) 6-mercaptopurine,
- l) cytosine arabinoside,
- m) podophyllotoxin,
 - n) etoposide,
 - o) etoposide phosphate,
 - p) melphalan,
 - q) vinblastine,
- r) vincristine,
 - s) leurosidine,
 - t) vindesine,
 - u) estramustine,
 - v) cisplatin,
 - w) cyclophosphamide, and
 - x) leurosine.

45. The conjugate according to Claim 12 wherein the cytotoxic agent is selected from doxorubicin and vinblastine or a cytotoxic derivative thereof.

5.16. The conjugate according to Claim 12 wherein the cytotoxic agent is doxorubicin or a cytotoxic derivative thereof.

The conjugate according to Claim 16 of the formula I:

X_L - oligopeptide - R

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wherein:

oligopeptide is an oligopeptide which is specifically recognized by
the free prostate specific antigen (PSA) and is capable of being
proteolytically cleaved by the enzymatic activity of the free prostate
specific antigen;

XL is absent or is an amino acid selected from:

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CH₃O

ОН

I

- a) phenylalanine,
- b) leucine,
- c) valine,
- d) isoleucine,
- e) (2-naphthyl)alanine,
- f) cyclohexylalanine,
- g) diphenylalanine,
- h) norvaline, and
- j) norleucine;

R is hydrogen or -(C=O)R¹; and

 R^1 is C_1 - C_6 -alkyl or aryl.

748. The conjugate according to Claim 17 wherein:

oligopeptide is an oligomer that comprises an amino acid sequence selected from:

- a) AsnLysIleSerTyrGln|Ser (SEQ.ID.NO.: 13),
 - b) LysIleSerTyrGln|Ser (SEQ.ID.NO.: 14),
 - c) GlyGluAsnGlyValGlnLysAspValSerGlnXaaSerIleTyr|SerGlnThrGlu(SEQ.ID.NO.: 15),
 - d) GlyLysGlyIleSerSerGlnTyr|SerAsnThrGluGluArgLeu (SEQ.ID.NO.: 2),
 - e) AsnLysIleSerTyrTyr|Ser (SEQ.ID.NO.: 127),
 - f) AsnLysAlaSerTyrGln|Ser (SEQ.ID.NO.: 128),
 - g) SerTyrGln|SerSer (SEQ.ID.NO.: 129), and
- h) hArgTyrGln|SerSer (SEQ.ID.NO.: 141);

wherein hArg is homoarginine and Xaa is any natural amino acid;

XL is absent or is an amino acid selected from:

- a)leucine,
- b) isoleucine, and
- d) valine; and

R is acetyl, pivaloyl or benzoyl.

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The conjugate according to Claim 16 which is selected from:

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CH₂OH CH₃O OH

OH

wherein X is:

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HaN— AsnLyslleSerTyrGlnSer-

H₂N— AsnLyslleSerTyrGlnSerSer-

H₂N— AsnLyslleSerTyrGlnSerSerSer -

H₂N— AsnLyslleSerTyrGlnSerSerSerThr

H₂N—AsnLyslleSerTyrGlnSerSerSerThrGlu

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(SEQ.ID.NO.: 13),

(SEQ.ID.NO.: 16),

(SEQ.ID.NO.: 17),

(SEQ.ID.NO.:10),

(SEQ.ID.NO.: 3),

 H_2N — AlaAsnLyslleSerTyrGlnSerSerSerThrGlu — C — (SEQ.ID.NO.: 11),

	Ο
	AcHN—AlaAsnLyslleSerTyrGlnSerSerSerThr—C— (SEQ.ID.NO.: 117),
5	AcHN—AlaAsnLyslleSerTyrGlnSerSerSerThrLeu-C— (SEQ.ID.NO.: 70),
	AcHN—AlaAsnLysAlaSerTyrGlnSerAlaSerThrLeu-C- (SEQ.ID.NO.: 118),
	AcHN—AlaAsnLysAlaSerTyrGlnSerAlaSerLeu-C- (SEQ.ID.NO.: 119),
10	AcHN—AlaAsnLysAlaSerTyrGlnSerSerSerLeu-C- (SEQ.ID.NO.: 120),
	O II AcHN—AlaAsnLysAlaSerTyrGlnSerSerLeu-C- (SEQ.ID.NO.: 121),
15	O II AcHN—SerTyrGlnSerSerSerLeu-C- (SEQ.ID.NO.: 144),
	AcHN— hArgTyrGlnSerSerSerLeu-C- (SEQ.ID.NO.: 145),
20	O II AcHN—LysTyrGlnSerSerSerLeu—C— (SEQ.ID.NO.: 124), or
	AcHN—LysTyrGlnSerSerNle —C— (SEQ.ID.NO.: 146).

20. The conjugate according to Claim 15 of the formula

П:

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OH

N

CO₂CH₃

OH

COCOCH₃

OH

CONH - oligopeptide - C - NH₂

wherein:

oligopeptide is an oligopeptide which is specifically recognized by the free prostate specific antigen (PSA) and is capable of being proteolytically cleaved by the enzymatic activity of the free prostate specific antigen.

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add B'/

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